

0032T\*425260

1 2223 70  
SEQ 15 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 08 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 09 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 10 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 11 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 12 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 13 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG  
SEQ 14 (1) MCHQQLVISWFSVLFLASPLVAIWELKDVVVELDWYDPAPGEMVVLTCDTPEEDGITWTLDDQSSEVLG

71 140  
SEQ 15 (71) SGKTLTIQVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 08 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 09 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 10 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 11 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 12 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 13 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF  
SEQ 14 (71) TGKTLTIHVKEFGDAGQYTCRKGGEVLSHLLHKKEDGIWSTDILKDQKEPKNKTFRLRCEAKNYSGRF

141 209  
SEQ 15 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 08 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 09 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 10 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 11 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 12 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 13 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE  
SEQ 14 (141) TCWWLTTISTDLTFSVKSSRGSSDPQGVTCGAATLSAERVGRDN-KEYEYSVECEQEDSACPAAEESLPIE

Fig. 1A

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SEQ 15	(280)	KSKREKKDRVFTDKTSATVICRKNASISVRAQDRYSSSSWSEWASVPCS--	328
SEQ 08	(276)	KSKREKKDRUFTDKTSATVICRKNAKIRVQARDRYSSFWSEWASVCS--	
SEQ 09	(281)	KNKREKK--LFDQTSAKVTCHKDAKIRVQARDRYSSSSWSEWASVPCS--	
SEQ 10	(281)	KNKREKK--LFDQTSAKVTCHKDAKIRVQARDRYSSSSWSEWASVPCS--	
SEQ 11	(281)	KNKREKK--LFDQTSAKVTCHKDAKIRVQARDRYHSSSWSEWASVPCS--	
SEQ 12	(276)	KNKREKK--LFDQTSAKVTCHKDAKIRVQARDRYSSSSWSDWASVSCG--	
SEQ 13	(276)	KNKREDR--LFDQTSAKVTCHKDAKIRVQARDRYSSSSWSEWASVSGV	
SEQ 14	(281)	KNKREKKDRLSVDKTSAKVMCHKDAKIRVQARDRYSSSSWSEWASVPCS--	

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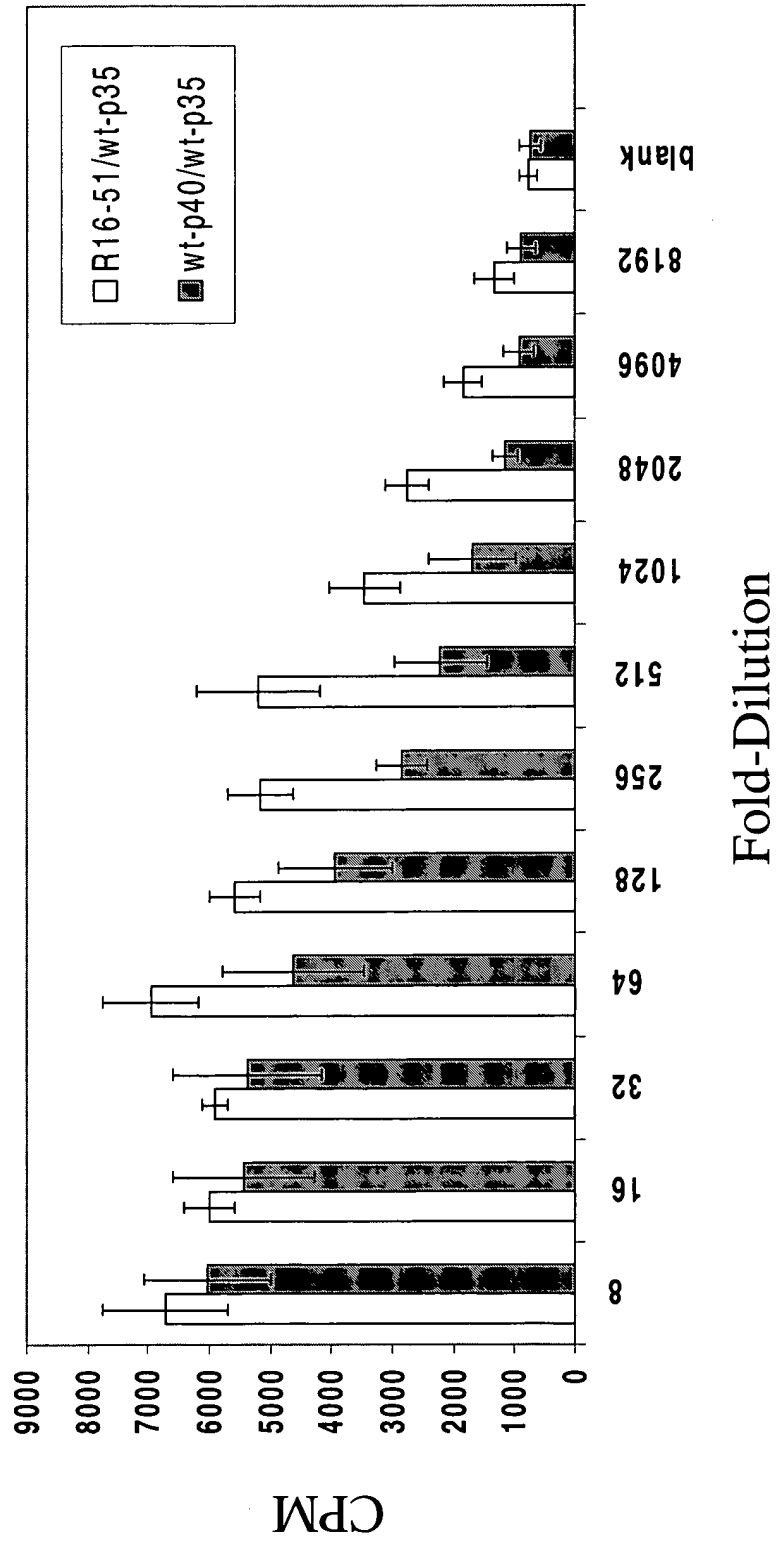


Fig. 2

003217-4232250

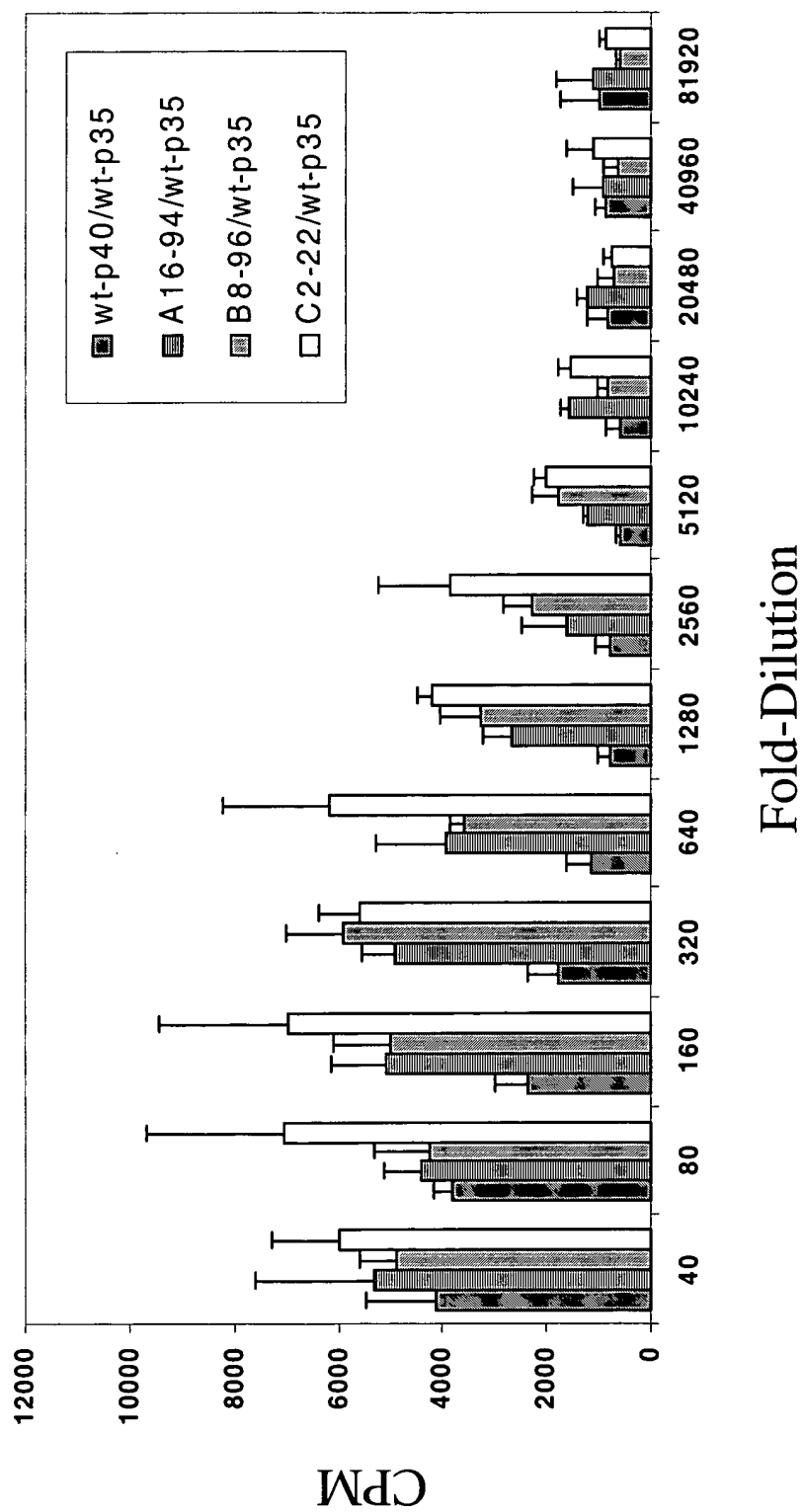


Fig. 3

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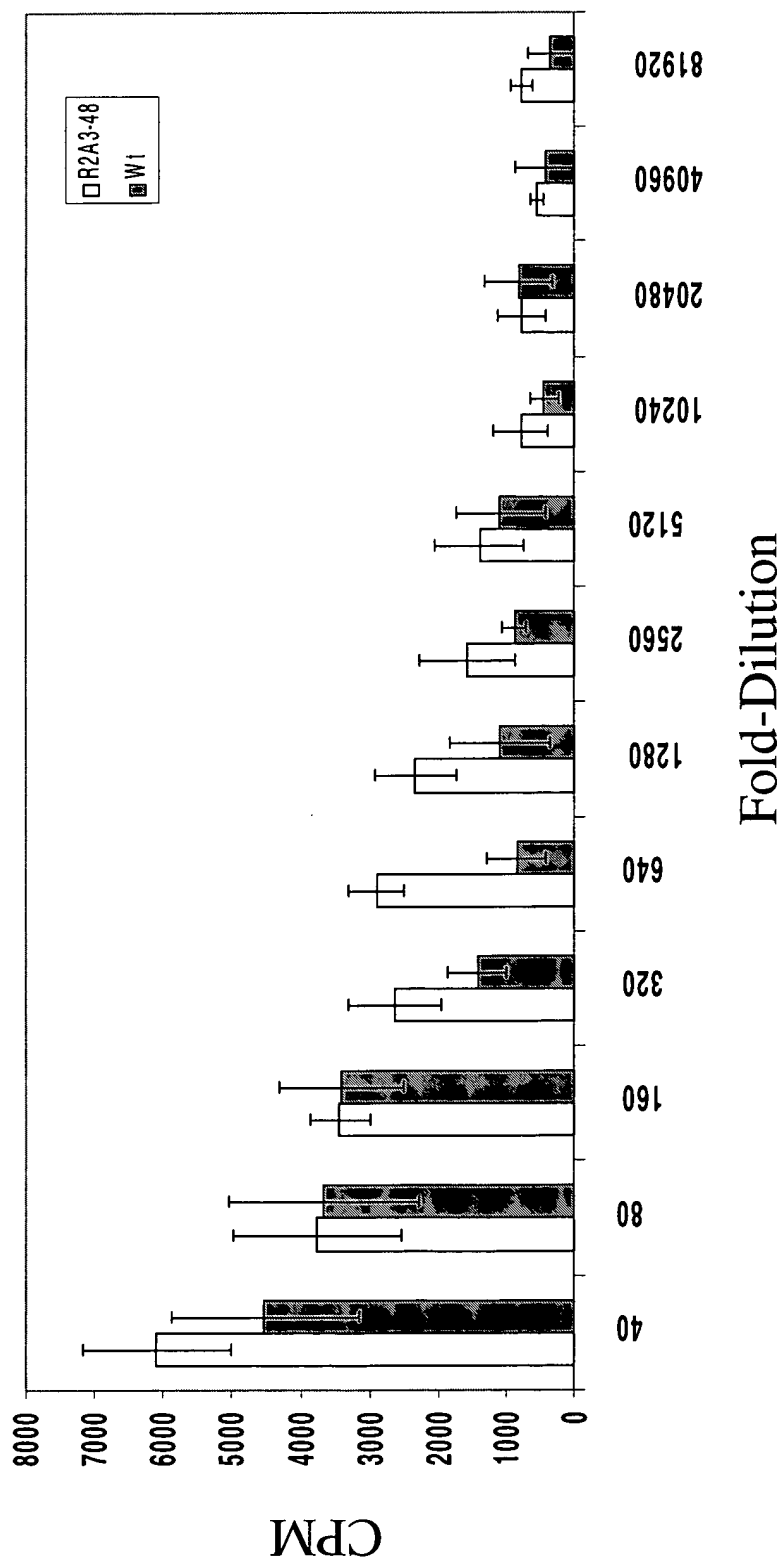
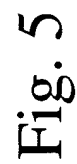


Fig. 4

[illegible]

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003277-4232250

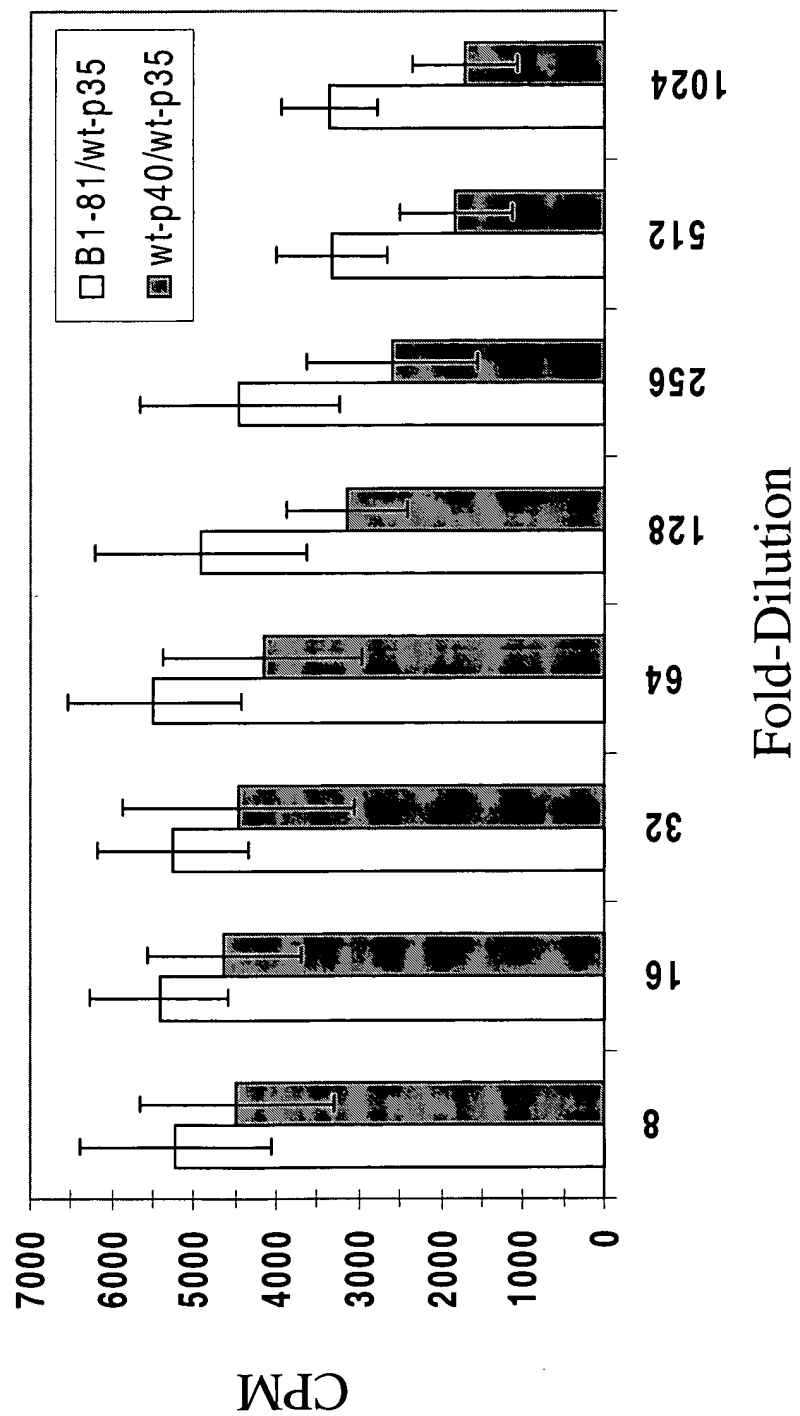
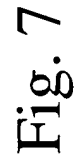


Fig. 6

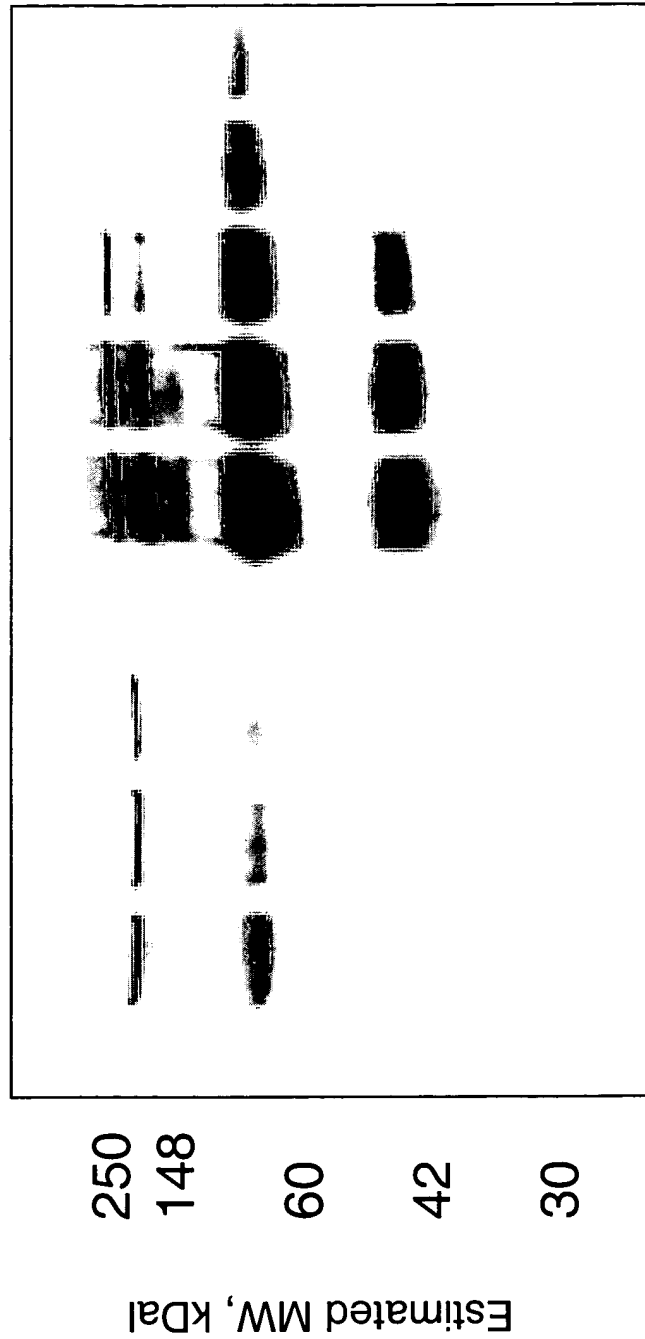
[illegible]



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wt-p40 / wt-p35

C2-22/wt-p35



Volume of protein solution

Fig. 8

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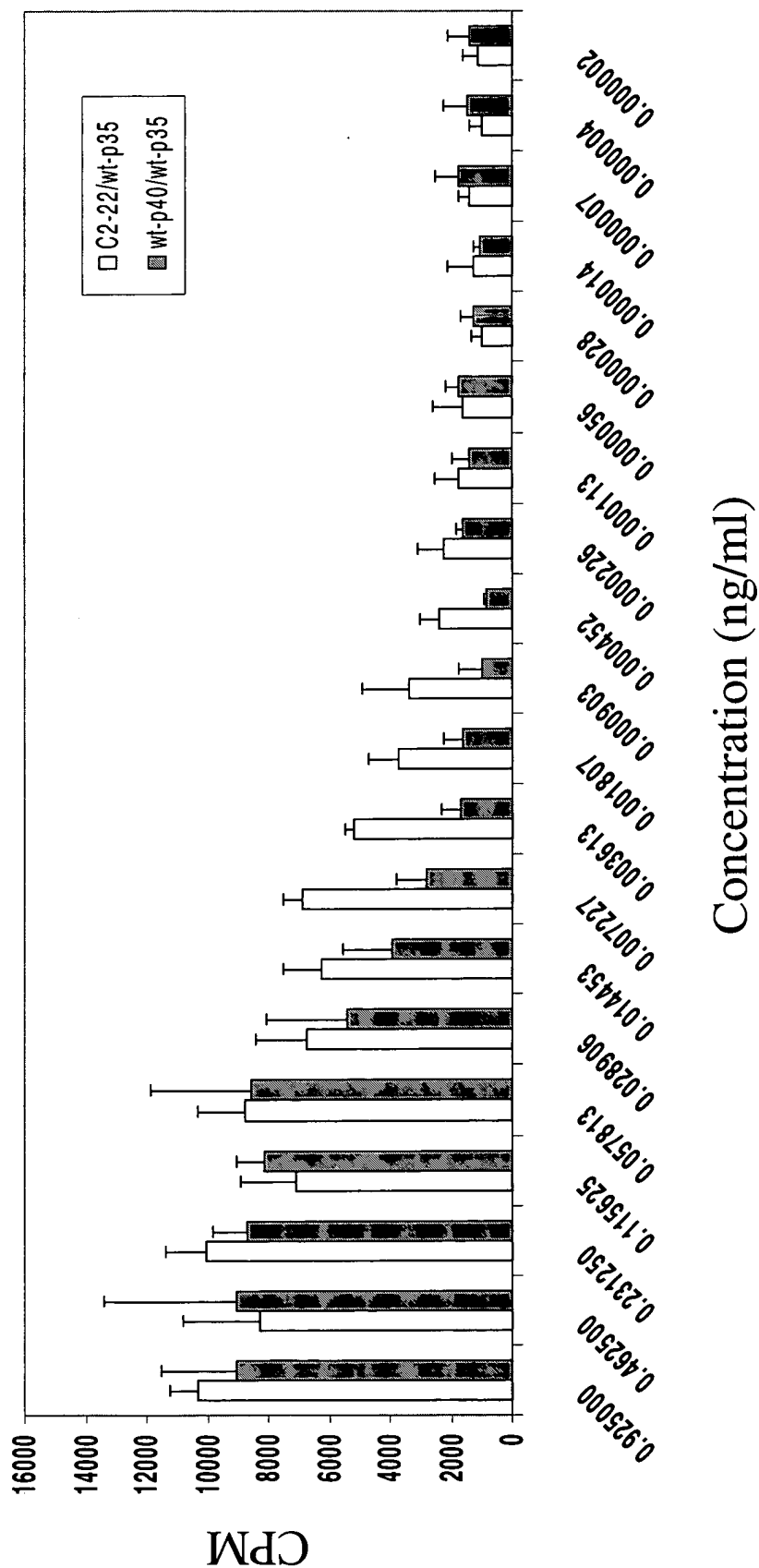


Fig. 9

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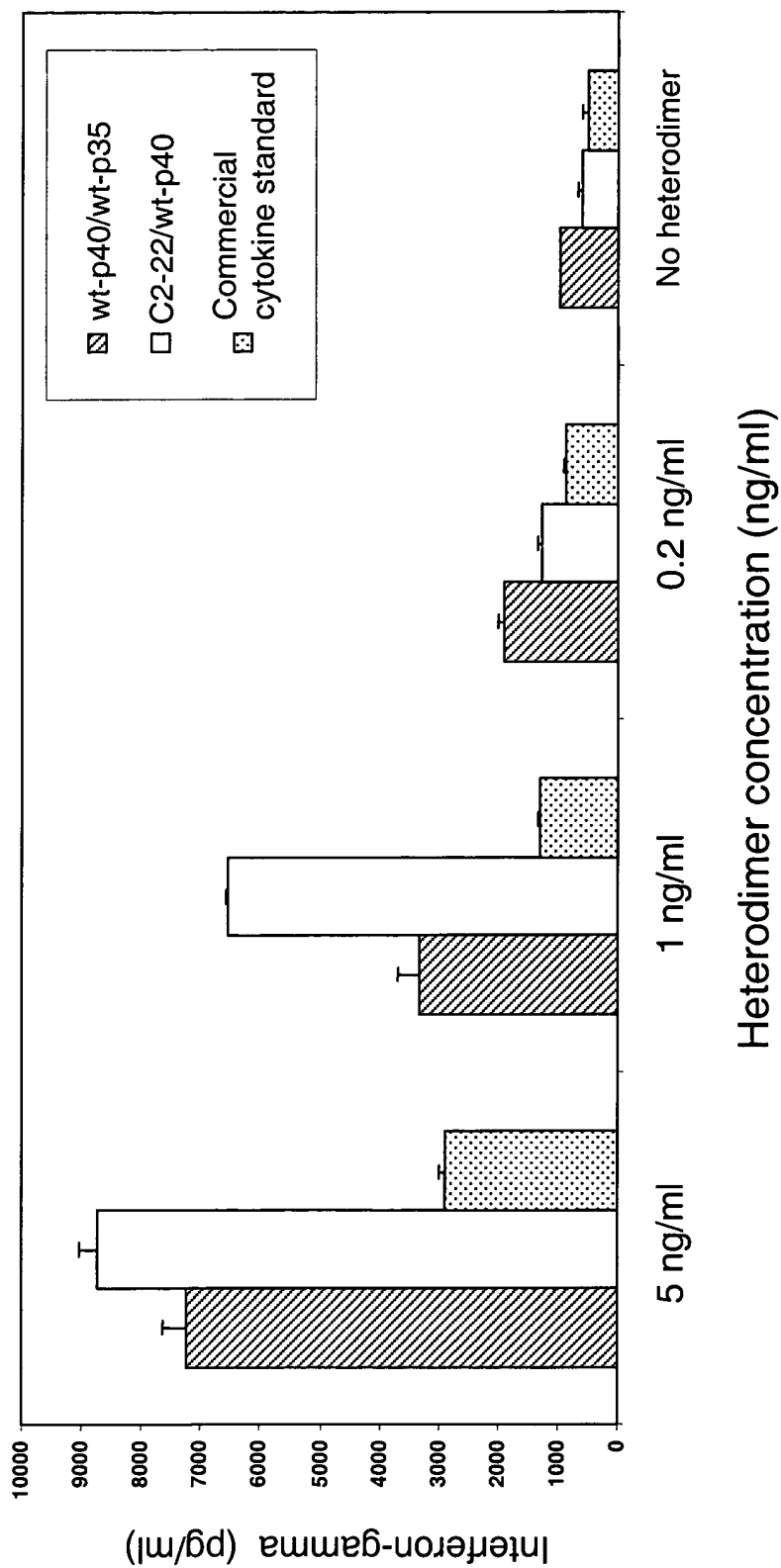


Fig. 10





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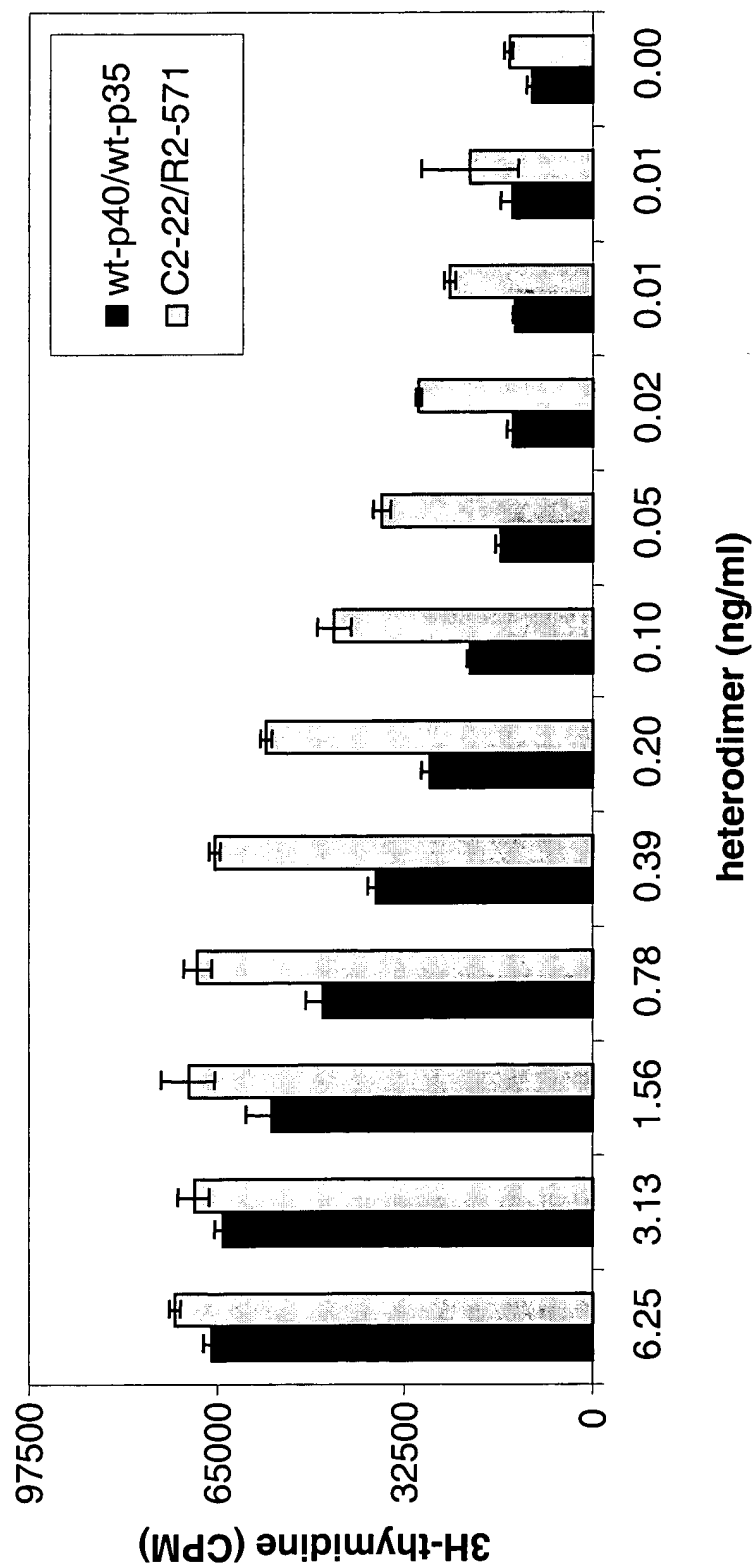


Fig. 12

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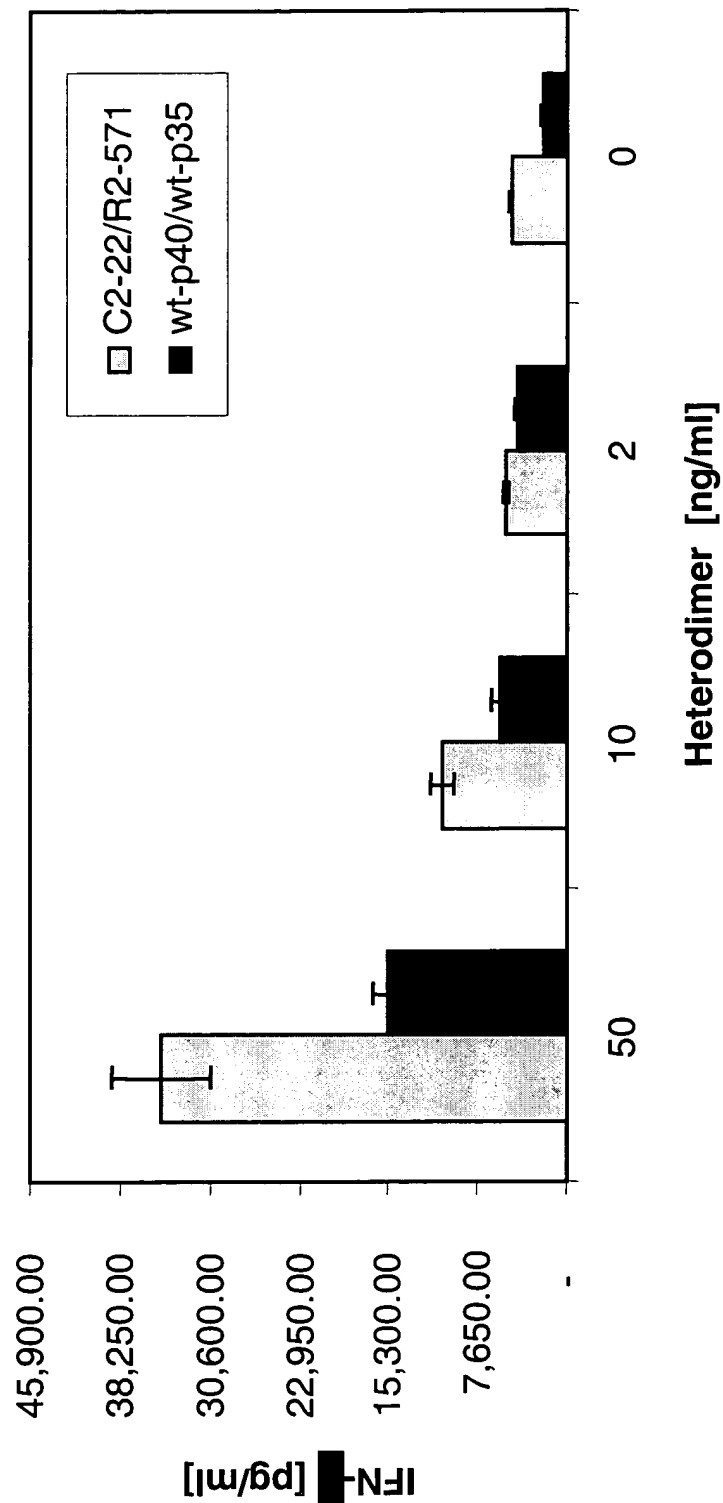


Fig. 13

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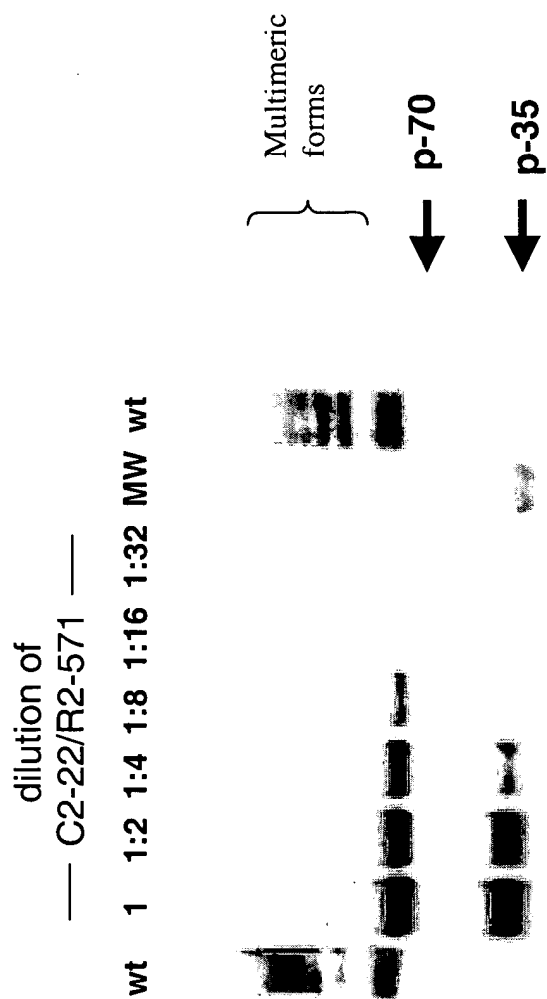


Fig. 14

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